

SEQUENCE LISTING

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Hirota, Kiyonori  
Sota, Hiroyuki

<120> Support having affinity for antibody

<130> 040894-7434-US

<140> 10575254  
<141> 2007-06-05

<150> US 10/575,254  
<151> 2006-04-10

<150> PCT/JP2004/014828  
<151> 2004-10-07

<150> JP 2003-352937  
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<170> PatentIn version 3.4

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Leu Asn Met Pro Asn Leu Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln  
20 25 30

Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
35 40 45

Lys Lys Leu Asn Glu Ser Gln Ala Pro Lys Gly Gly Gly Cys Ala  
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Asp Asp Asp Asp Asp Asp  
65 70

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Ala Asp Asn Asn Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile  
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Leu Asn Met Pro Asn Leu Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln  
20 25 30

Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ser Glu Ala  
35 40 45

Lys Lys Leu Asn Glu Ser Gln Ala Pro Lys Ala Asp Asn Asn Phe Asn  
50 55 60

Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu Asn Met Pro Asn Leu  
65 70 75 80

Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu Lys Asp Asp Pro  
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Ser Gln Ser Ala Asn Leu Leu Ser Glu Ala Lys Lys Leu Asn Glu Ser  
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Gln Ala Pro Lys Gly Gly Gly Cys Ala Asp Asp Asp Asp Asp Asp  
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Ala Asp Asn Asn Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile  
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Leu Asn Met Pro Asn Leu Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln  
20 25 30

Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
35 40 45

Lys Lys Leu Asn Glu Ser Gln Ala Pro Lys  
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<223> A domain dimer

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Leu Asn Met Pro Asn Leu Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln

20 25 30

Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ser Glu Ala

35 40 45

Lys Lys Leu Asn Glu Ser Gln Ala Pro Lys Ala Asp Asn Asn Phe Asn

50 55 60

Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu Asn Met Pro Asn Leu

65 70 75 80

Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu Lys Asp Asp Pro

85 90 95

Ser Gln Ser Ala Asn Leu Leu Ser Glu Ala Lys Lys Leu Asn Glu Ser

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Gln Ala Pro Lys Gly Gly Gly Cys Ala Asp Asp Asp Asp Asp Asp

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<212> DNA

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<223> DNA encoding protein for antibody immobilization

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<223> DNA encoding protein for antibody immobilization

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caaagtgcta acctattgtc agaagctaaa aagttaaatg aatctcaagc accgaaaggt 180  
gataacaatt tcaacaaaga acaacaaaat gctttctatg aaatcttgaa tatgcctaac 240  
ttaaacaacgaa aacaacgcaa tggtttcatc caaagcttaa aagatgaccc aagccaaagt 300  
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tctatgaaat ttcatatgtt cctaacttaa acgaagaaca acgcaatggt ttcatccaaa 180  
gcttaaaaaga tgacccaagc caaagtgcta acctattgtc agaagctaaa aagttaaatg 240  
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tctatgaaat ttcatatgtt cctaacttaa acgaagaaca acgcaatggt ttcatccaaa 180  
gcttaaaaaga tgacccaagc caaagtgcta acctattgtc agaagctaaa aagttaaatg 240  
aatctcaagc accgaaaggt gataacaatt tcaacaaaga acaacaaaat gctttctatg 300

aaatcttcaa tatgcctaac taaaacgaag aacaacgcaa tggtttcatc caaagcttaa 360  
aagatgaccc aagccaaagt gctaacctat tgtcagaagc taaaaagtta aatgaatctc 420  
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